Donut Analysis with Root

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Outline

- Root
- Donut analysis
- Root and Donut
- Emulsion data tree
- Histograms, tracks
- Magnetic moment analysis
- Result
- Root outlook

Root introduction

- Data analysis framework
 - Object oriented
 - C++ based
- developed at CERN by Rene Brun, Fons Rademakers
- many people, experiments are now using root
 - NA49, ATLAS, RHIC, STAR, ALICE, HADES,...
- Root is the successor to PAW, Geant, CERNLIB

Donut analysis (to date)

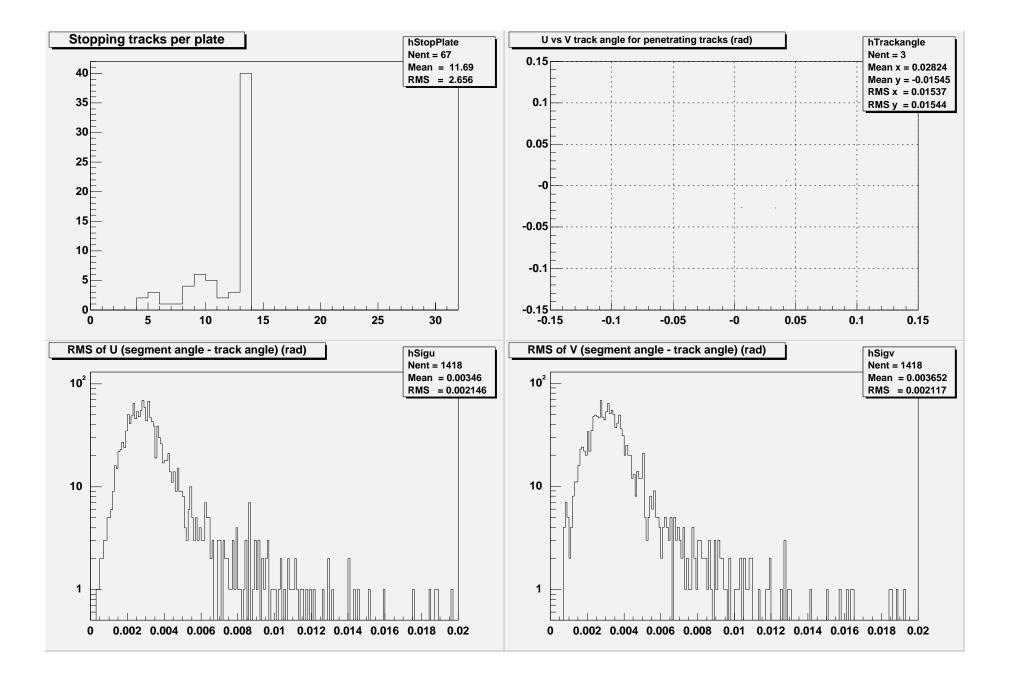
- Spectrometer data is analyzed with Fortran/Cernlib based software
- Emulsion data is analyzed in Nagoya
 - alignment in a C program (Aoki)
 - vertex finding:
 - in a simple C program (Aoki)
 - visually (Okada, using software based on Aokis C code)
 - in a Fortran program (Komatsu)
 - also here in the US in connection with the spectrometer (Bruce)

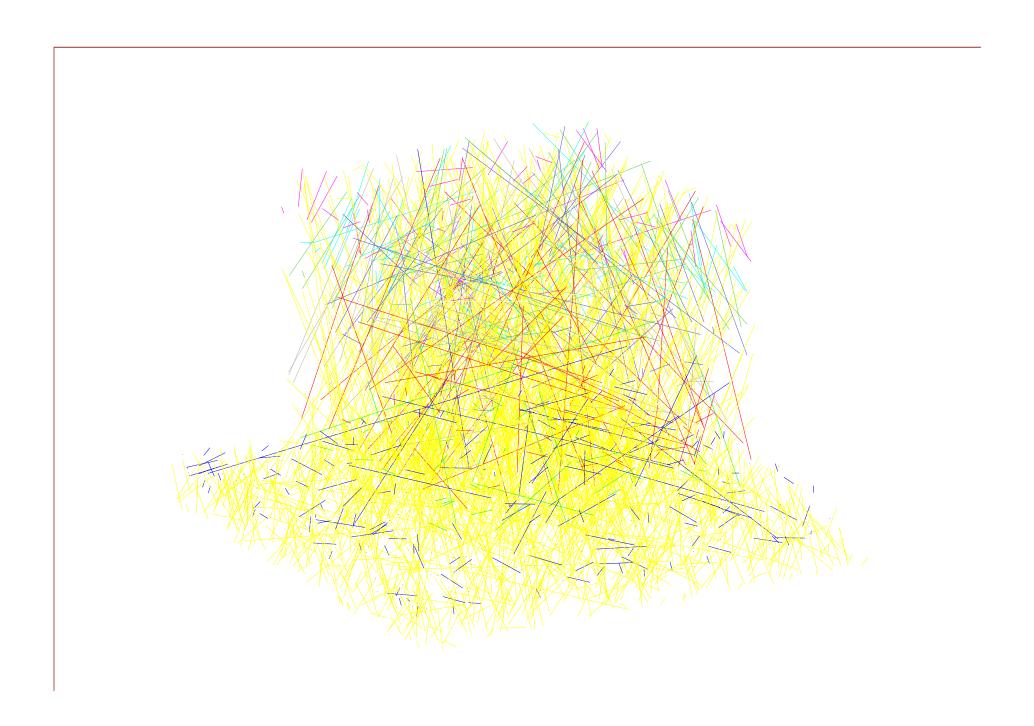
Root and Donut

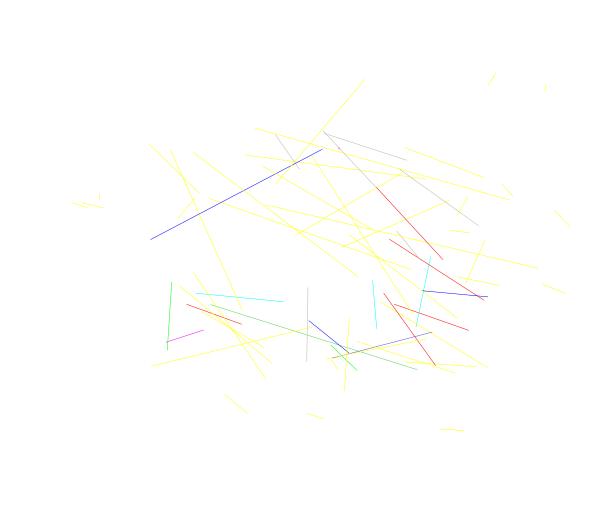
- Emulsion data: OO
 - Emulsion track segments
 - Emulsion tracks
 - Vertices
 - (Spectrometer tracks)
- We are starting to analyze emulsion data
 - alignment?
 - Vertex finding
 - decay search
 - magnetic moment analysis
- Some analysis tools (routines) to do this work exist
 - there are different tools for different tasks
 - there are no tools do decay search or my μ_{ν} analysis

Donut Root Tree

- I started to put emulsion information into a root framework
- The "Tree" has branches:
 - event information (run, event number, scan volume, ...)
 - emulsion track segment information
 - emulsion track information
 - spectrometer track information
 - emulsion vertex information
 - 4 histograms (stopping tracks/plate; U,V angle alignment; track angle)

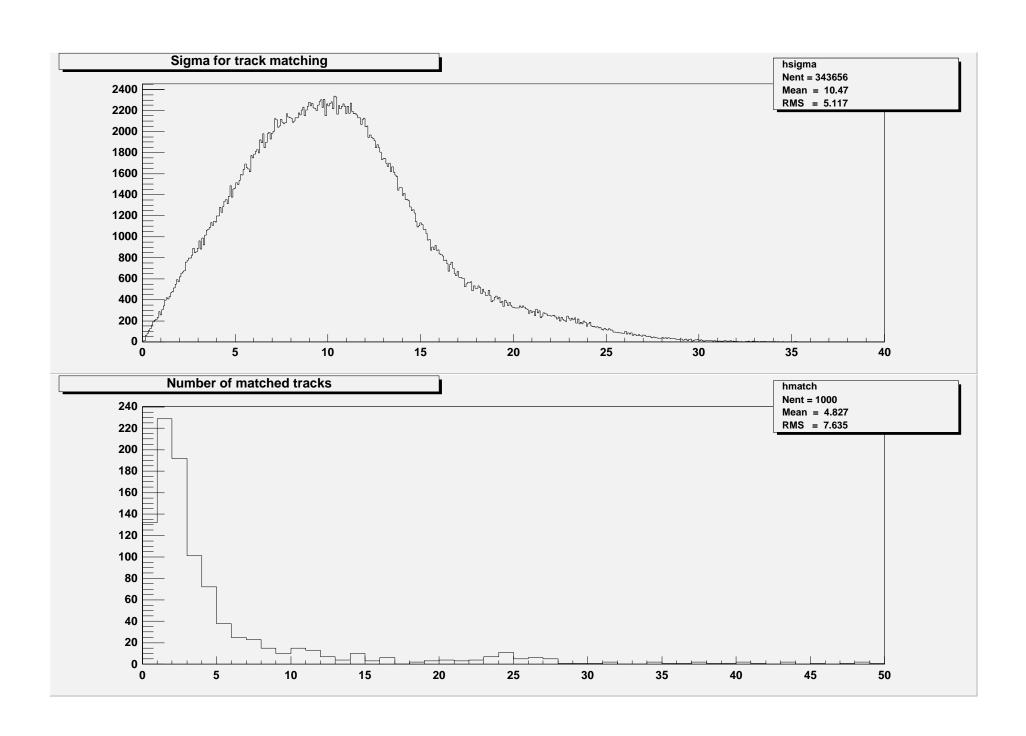


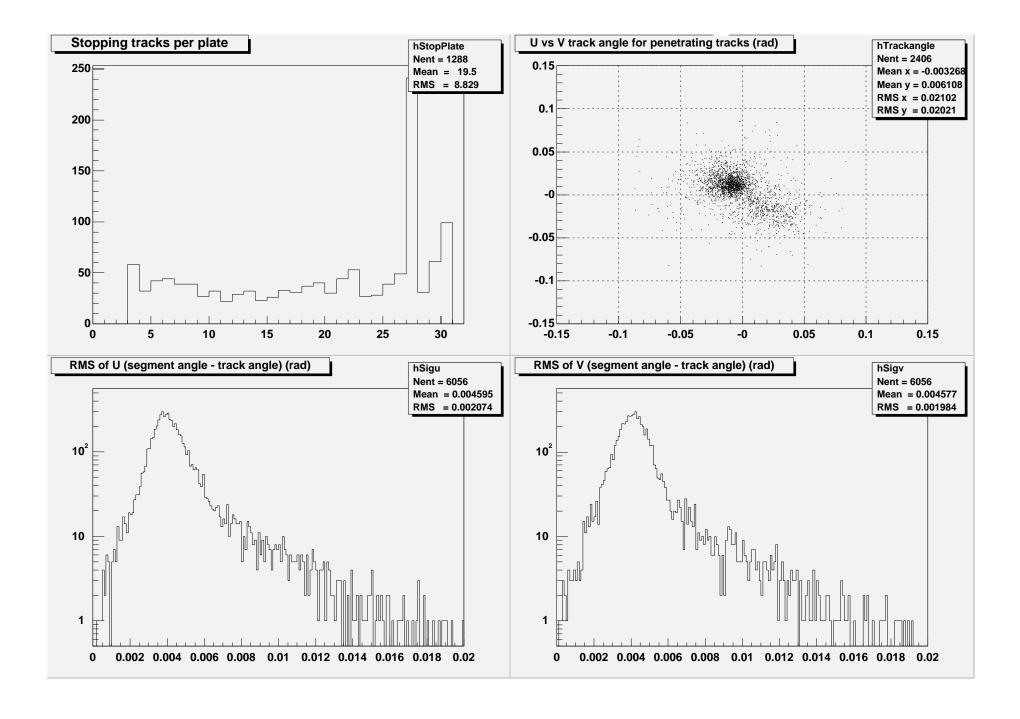




Magnetic moment analysis

- My candidate event sample contains events with single tracks
- I prepared predictions for a scanning volume
- The events were scanned in Nagoya
- Analysis:
 - match spectrometer track to stopping tracks
 - 90% CL volume in u,v,θu, θv space
 - check: create a random track and match to stopping tracks





Result

- The candidate event has 94 stopping-andexiting track matches to the spectrometer track
- the single track is in the high-muon-flux region
- I need
 - better alignment (reduce # of stopping tracks)
 - refined analysis

Root outlook

- The tree has been created
- A simple analysis is possible
- More work needs to be done
 - vertex branch
 - better implementation of spectrometer tracks
 - analysis routines
 - Nagoya routines
- I would like to encourage people to use it for emulsion analysis